RAW SEQUENCE LISTING DATE: 02/04/2003
PATENT APPLICATION: **US/10/007,621D** TIME: 13:47:20

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\02042003\J007621D.raw**

```
      3 <110> APPLICANT: Visigen Biotechnologies, Inc.
      5 <120> TITLE OF INVENTION: Enzymatic Nucleic Acid Synthesis: Compositions and Methods
for Altering
      6       Monomer Incorporation Fidelity
      8 <130> FILE REFERENCE: 00007/02PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/007,621D
     11 <141> CURRENT FILING DATE: 2001-12-03
     13 <150> PRIOR APPLICATION NUMBER: 60/250,764
     14 <151> PRIOR FILING DATE: 2000-12-01
     16 <160> NUMBER OF SEQ ID NOS: 9
     18 <170> SOFTWARE: PatentIn version 3.1
     20 <210> SEQ ID NO: 1
     21 <211> LENGTH: 7
     22 <212> TYPE: DNA
     23 <213> ORGANISM: Artificial
     25 <220> FEATURE:
     26 <223> OTHER INFORMATION: The sequences listed here are artifically generated DNA
sequences
     27       synthesized to test fidelity of monomer incorporation due to sub
     28       stitution at the gamma phosphate of the dNTPs.
     30 <220> FEATURE:
W--> 31 <221> NAME/KEY: Oligonucleotide
     32 <222> LOCATION: (1)..(7)
     33 <223> OTHER INFORMATION: An example of an oligonucleotide discussed the in the
definition
     34       section of the application.
     37 <400> SEQUENCE: 1
     38 atgcctg                                                              7
     41 <210> SEQ ID NO: 2
     42 <211> LENGTH: 19
     43 <212> TYPE: DNA
     44 <213> ORGANISM: Artificial
     46 <220> FEATURE:
     47 <223> OTHER INFORMATION: This sequence is a primer strand for Taq DNA polymerase.
     49 <220> FEATURE:
     50 <221> NAME/KEY: primer_bind
     51 <222> LOCATION: (1)..(19)
     52 <223> OTHER INFORMATION: Primer strand for Taq DNA polymerase
     55 <400> SEQUENCE: 2
     56 ggtactaagc ggccgcatg                                                19
     59 <210> SEQ ID NO: 3
     60 <211> LENGTH: 20
     61 <212> TYPE: DNA
     62 <213> ORGANISM: Artificial
```

64 <220> FEATURE:
65 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence 2 wi

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```
     66       th the addition of a T residue at the end of the strand designate
     67       d BOT-T 3'.
     69 <220> FEATURE:
W--> 70 <221> NAME/KEY: Template
     71 <222> LOCATION: (1)..(19)
     72 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
     75 <400> SEQUENCE: 3
     76 ccatgattcg ccggcgtact                                          20
     79 <210> SEQ ID NO: 4
     80 <211> LENGTH: 20
     81 <212> TYPE: DNA
     82 <213> ORGANISM: Artificial
     84 <220> FEATURE:
     85 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence
2 wi
     86       th the addition of a C residue at the end of the strand designate
     87       d BOT-C 3'.
     89 <220> FEATURE:
W--> 90 <221> NAME/KEY: Template
     91 <222> LOCATION: (1)..(19)
     92 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
     95 <400> SEQUENCE: 4
     96 ccatgattcg ccggcgtacc                                          20
     99 <210> SEQ ID NO: 5
    100 <211> LENGTH: 20
    101 <212> TYPE: DNA
    102 <213> ORGANISM: Artificial
    104 <220> FEATURE:
    105 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence
2 wi
    106       th the addition of a G residue at the end of the strand designate
    107       d BOT-G 3'.
    109 <220> FEATURE:
W--> 110 <221> NAME/KEY: Template
    111 <222> LOCATION: (1)..(19)
    112 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
    115 <400> SEQUENCE: 5
    116 ccatgattcg ccggcgracg                                          20
    119 <210> SEQ ID NO: 6
    120 <211> LENGTH: 20
    121 <212> TYPE: DNA
    122 <213> ORGANISM: Artificial
    124 <220> FEATURE:
    125 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence
2 wi
    126       th the addition of a A residue at the end of the strand designate
    127       d BOT-A 3'.
    129 <220> FEATURE:
W--> 130 <221> NAME/KEY: Template
    131 <222> LOCATION: (1)..(19)
    132 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
    135 <400> SEQUENCE: 6
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136 ccatgattcg ccggcgtaca 20
139 <210> SEQ ID NO: 7
140 <211> LENGTH: 23
141 <212> TYPE: DNA
142 <213> ORGANISM: Artificial
144 <220> FEATURE:
145 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence 2 wi
146 th the addition of a TAG residues at the end of the strand design
147 ated BOT-Sau 3'.
149 <220> FEATURE:
W--> 150 <221> NAME/KEY: Template
151 <222> LOCATION: (1)..(19)
152 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
155 <400> SEQUENCE: 7
156 ccatgattcg ccggcgtacc tag 23
159 <210> SEQ ID NO: 8
160 <211> LENGTH: 21
161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial
164 <220> FEATURE:
165 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence 2 wi
166 th the addition of a TC residues at the end of the strand designa
167 ted BOT-TC 3'.
169 <220> FEATURE:
W--> 170 <221> NAME/KEY: Template
171 <222> LOCATION: (1)..(19)
172 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
175 <400> SEQUENCE: 8
176 ccatgattcg ccggcgtact c 21
179 <210> SEQ ID NO: 9
180 <211> LENGTH: 23
181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Template Strand - antisense to the primer strand of sequence 2 wi
186 th the addition of a TTTC residues at the end of the strand desig
187 nated BOT-3TC 3'.
189 <220> FEATURE:
W--> 190 <221> NAME/KEY: Template
191 <222> LOCATION: (1)..(19)
192 <223> OTHER INFORMATION: Anti-sense to the primer sequence 2.
195 <400> SEQUENCE: 9
196 ccatgattcg ccggcgtact ttc 23

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/04/2003
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Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9

VERIFICATION SUMMARY DATE: 02/04/2003
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Input Set : **A:\EP.txt**
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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:31 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:70 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:90 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:110 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:130 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:150 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:170 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:190 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9